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APPLICATION NO.	FILING DATE	FIRST NAMED INVENTOR	ATTORNEY DOCKET NO.	CONFIRMATION NO.
09/847,539	05/01/2001	Lars Bjorck	100084.415US	4148

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David Kalow  
Kalow & Springut LLP  
488 Madison Avenue 19th Floor  
New York, NY 10022

EXAMINER

BASKAR, PADMAVATHI

ART UNIT

PAPER NUMBER

1645

DATE MAILED: 09/24/2003

19

Please find below and/or attached an Office communication concerning this application or proceeding.

<b>Office Action Summary</b>	Application No.	Applicant(s)
	09/847,539	BJORCK ET AL.
	Examiner Padmavathi v Baskar	Art Unit 1645
-- The MAILING DATE of this communication appears on the cover sheet with the correspondence address --		
<b>Period for Reply</b>		
A SHORTENED STATUTORY PERIOD FOR REPLY IS SET TO EXPIRE <u>3</u> MONTH(S) FROM <b>THE MAILING DATE OF THIS COMMUNICATION.</b>		
<ul style="list-style-type: none"> <li>- Extensions of time may be available under the provisions of 37 CFR 1.136(a). In no event, however, may a reply be timely filed after SIX (6) MONTHS from the mailing date of this communication.</li> <li>- If the period for reply specified above is less than thirty (30) days, a reply within the statutory minimum of thirty (30) days will be considered timely.</li> <li>- If NO period for reply is specified above, the maximum statutory period will apply and will expire SIX (6) MONTHS from the mailing date of this communication.</li> <li>- Failure to reply within the set or extended period for reply will, by statute, cause the application to become ABANDONED (35 U.S.C. § 133).</li> <li>- Any reply received by the Office later than three months after the mailing date of this communication, even if timely filed, may reduce any earned patent term adjustment. See 37 CFR 1.704(b).</li> </ul>		
<b>Status</b>		
1) <input checked="" type="checkbox"/> Responsive to communication(s) filed on <u>23 July 2003</u> .		
2a) <input checked="" type="checkbox"/> This action is <b>FINAL</b> .                            2b) <input type="checkbox"/> This action is non-final.		
3) <input type="checkbox"/> Since this application is in condition for allowance except for formal matters, prosecution as to the merits is closed in accordance with the practice under <i>Ex parte Quayle</i> , 1935 C.D. 11, 453 O.G. 213.		
<b>Disposition of Claims</b>		
4) <input checked="" type="checkbox"/> Claim(s) <u>23 and 25 – 38</u> is/are pending in the application.		
4a) Of the above claim(s) _____ is/are withdrawn from consideration.		
5) <input type="checkbox"/> Claim(s) _____ is/are allowed.		
6) <input checked="" type="checkbox"/> Claim(s) <u>23 and 25 – 38</u> is/are rejected.		
7) <input type="checkbox"/> Claim(s) _____ is/are objected to.		
8) <input type="checkbox"/> Claim(s) _____ are subject to restriction and/or election requirement.		
<b>Application Papers</b>		
9) <input type="checkbox"/> The specification is objected to by the Examiner.		
10) <input type="checkbox"/> The drawing(s) filed on _____ is/are: a) <input type="checkbox"/> accepted or b) <input type="checkbox"/> objected to by the Examiner. Applicant may not request that any objection to the drawing(s) be held in abeyance. See 37 CFR 1.85(a).		
11) <input type="checkbox"/> The proposed drawing correction filed on _____ is: a) <input type="checkbox"/> approved b) <input type="checkbox"/> disapproved by the Examiner. If approved, corrected drawings are required in reply to this Office action.		
12) <input type="checkbox"/> The oath or declaration is objected to by the Examiner.		
<b>Priority under 35 U.S.C. §§ 119 and 120</b>		
13) <input type="checkbox"/> Acknowledgment is made of a claim for foreign priority under 35 U.S.C. § 119(a)-(d) or (f). a) <input type="checkbox"/> All b) <input type="checkbox"/> Some * c) <input type="checkbox"/> None of: 1. <input type="checkbox"/> Certified copies of the priority documents have been received. 2. <input type="checkbox"/> Certified copies of the priority documents have been received in Application No. _____. 3. <input type="checkbox"/> Copies of the certified copies of the priority documents have been received in this National Stage application from the International Bureau (PCT Rule 17.2(a)). * See the attached detailed Office action for a list of the certified copies not received.		
14) <input type="checkbox"/> Acknowledgment is made of a claim for domestic priority under 35 U.S.C. § 119(e) (to a provisional application). a) <input type="checkbox"/> The translation of the foreign language provisional application has been received.		
15) <input type="checkbox"/> Acknowledgment is made of a claim for domestic priority under 35 U.S.C. §§ 120 and/or 121.		
<b>Attachment(s)</b>		
1) <input type="checkbox"/> Notice of References Cited (PTO-892)		
2) <input type="checkbox"/> Notice of Draftsperson's Patent Drawing Review (PTO-948)		
3) <input type="checkbox"/> Information Disclosure Statement(s) (PTO-1449) Paper No(s) _____.		
4) <input type="checkbox"/> Interview Summary (PTO-413) Paper No(s) _____.		
5) <input type="checkbox"/> Notice of Informal Patent Application (PTO-152)		
6) <input type="checkbox"/> Other: _____.		

***Response to Amendment***

1. Applicant's amendment filed on 6/23/03 (Paper # 17) is acknowledged. Claims 24 and 39 have been canceled. Claims 23, 25 and 29 ~~have~~ been amended. Claims 23, 25-38 are under examination.

***Priority***

2. Acknowledgment is made of applicant's claim for foreign priority under 35 U.S.C. 119(a)(d) to application United Kingdom 9823975.9. The certified copy of United Kingdom 9823975.9 has been received and foreign priority has been accorded as of the filing date of the priority document, 11/2/1998.
3. The examiner acknowledges the new title and has been placed in the application.

***Rejection Withdrawn***

4. In view of amendment to the claims, the rejection of claims 23 and 25 -38 under 35 U.S.C. 101 non-statutory subject matter is withdrawn.

***Rejections Maintained***

5. The rejection of claims 23, 25-38 under 35 U.S.C. 112, first paragraph, because the specification, while being enabling for an isolated protein that is capable of binding to  $\alpha_2$  M that comprises the amino acid sequence of SEQ.ID.NO: 6 does not reasonably provide enablement for a functional variant thereof that is capable of binding to  $\alpha_2$  M and has at least 78% homology to amino acids to SEQ.ID.NO: 6 over at least 30 amino acids or over at least 100 amino acids or a variant with at least 75% homology to amino acids 59 to 86 of SEQ.ID.NO: 6 or a peptide fragment comprising 6, 15 and 20 amino acids of SEQ.ID.NO: 6 or variants with at least 75%

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and 78% homology to amino acids 59 to 86 of SEQ.ID.NO: 6 or functional variant comprising one or more tandem repeats is maintained as set forth in the previous office action.

The specification is not enabled for a functional variant thereof or functional variant with 75% and 78% homology to amino acids 59 to 86 of SEQ.ID.NO: 6 or peptide fragments comprising 6, 15 and 20 amino acids of SEQ.ID.NO: 6 or variants with at least 75% and 78% homology to amino acids 59 to 86 of SEQ.ID.NO: 6 because the specification is totally silent. If it is unclear to one skilled in the art what are those functional equivalents or sequences are embraced by the claim since the specification lacks the algorithm and parameters used to determine percent identity or derivatives of fragments.

The specification is silent in what changes have been made to SEQ.ID.NO: 6 to obtain the above functional variants. It is well known that for proteins, for example, even a single amino acid change can destroy the function of the biomolecule. The effects of these changes are largely unpredictable as to which ones have a significant effect versus not. Further, specification is silent on how to make these proteins with sequence homology or variants or fragments. What changes would have an adverse effect on the function of this peptide is not predictable. It is known in the art that derivatives or variants, which are obtained by substitutions, deletions, or modifications of the amino acids of a protein, alter the function of the protein. The amino acid sequence of a protein determines its structural and functional properties, predictability of which changes can be tolerated in a protein's amino acid sequence and still retain similar activity requires a knowledge of and guidance with regard to which amino acids in the protein's sequence, if any, are tolerant of modification and which are conserved (i.e. expected intolerant to modification), and detailed knowledge of the ways in which the proteins' structure relates to its function. However, the problem of predicting protein structure from mere sequence data of a single protein and in turn utilizing predicted structural determinations to ascertain functional aspects of the protein and finally what changes can be tolerated with respect thereto is extremely complex (Bowie et al. Science, Vol. 247: 1990; p. 1306; p. 1308) and is well outside the realm of routine experimentation.

Applicant's arguments filed on 6/23/03 have been fully considered but they are not deemed to be persuasive.

Applicant states that in light of the amendments made to the claims, the rejection under 35 U.S.C. 112, first paragraph is inappropriate, variant as described in page 6, lines 4-6 can be used in vaccine formulation for generating preferably a protective immune response, procedures identifying candidates for generating an immune response have been illustrated in page 11, lines 14-21 and pages 7 and 14 teach calculating homology and amino acid identity and therefore, the rejection should be withdrawn.

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The examiner has reviewed the citations which applicant pointed to the examiner and found no support for the claimed language. While specification teaches general methods of calculating homology but does not teach how to make and use the variants as claimed in the specification. The variants, which bind to  $\alpha_2$  macroglobulin, have not been taught by the specification. The state of the art teaches (see IDS, AQ reference, first two paragraphs) that the interaction between streptococci and the two forms of  $\alpha_2$  macroglobulin ( $\alpha_2$  M) is highly specific. While human pathogenic streptococci bind to native form of  $\alpha_2$  M, bovine and equine streptococci interact with proteinase complexed form of  $\alpha_2$  M. Therefore, the amino acid sequence of a protein determines its structural and functional properties such as binding to the  $\alpha_2$  M. What changes can be tolerated in a protein to retain its functional activity requires a knowledge of and guidance with regard to which amino acids in the protein's sequence, if any, are tolerant of modification and which are conserved, and detailed knowledge of the ways in which the proteins' structure relates to its function have not been disclosed with regard to variants. However, the problem of predicting protein structure from mere sequence data of a single protein and in turn utilizing predicted structural determinations to ascertain functional aspects of the protein and finally what changes can be tolerated with respect thereto is extremely complex (Bowie et al. Science, Vol. 247: 1990; p. 1306; p. 1308) and is well outside the realm of routine experimentation.

7. The rejection of claims 23 and 25 – 38 under 35 U.S.C. 102(b) as being anticipated by Lammler et al 1986 (Zbl.Bakt.Hyg A 261, 161-166) is maintained as set forth in the previous office action.

Lammler et al disclose ten of the 11 streptococcal cultures of serological group A, T type 4 possessing antigen (T4) bound to  $\alpha_2$  macroglobulin (see abstract, Table 1). Further, the prior art disclose that only one strain 71-715 did not bind to  $\alpha_2$  macroglobulin but other 10 strains including S.pyogenes strain 71-727 bound to  $\alpha_2$  macroglobulin (see page 163 under binding of

plasma proteins). Thus the prior art reads on claim 28 as well. In the absence of evidence to the contrary the disclosed prior art streptococcal proteins comprise amino acid sequence of SEQ.ID.NO: 6. Since the Office does not have the facilities for examining and comparing applicants' claimed protein with the T4 antigen of the prior art, the burden is on applicant to show a novel or unobvious difference between the claimed protein and the T 4 antigen of the prior art. See *In re Best*, 562 F.2d 1252, 195 USPQ 430 (CCPA 1977).

Applicant's arguments filed on 6/23/03 have been fully considered but they are not deemed to be persuasive.

Applicant states that the cited reference did not disclose the sequence, SEQ.ID.NO: 6 or variant.

The examiner has rejected the claims based on inherency. Protein T4 (antigen) bound to the  $\alpha_2$  macroglobulin and therefore, it contains the amino acid sequence SEQ.ID.NO: 6. Applicant did not show any evidence that the disclosed protein and the claimed invention are not the same.

8. The rejection of claims 23 and 25 – 38 under 35 U.S.C. 102(b) as being anticipated by Chhatwal et al (J.Bacteriology, 169; 3691-3695) is maintained as set forth in the previous office action.

Chhatwal et al disclose streptococcal cultures of serological group A, including *S.pyogenes* strain A8189 bound to  $\alpha_2$  macroglobulin (see abstract, Table 1 and figure 3). Further, the prior art disclose streptococcal lysates and purified proteins bind to  $\alpha_2$  macroglobulin as well as antibodies to  $\alpha_2$  macroglobulin indicating that the protein binds to  $\alpha_2$  macroglobulin (see figure 4). *S.pyogenes* strain A8189 bound to  $\alpha_2$  macroglobulin (figures 1-4) since this strain is not SF 370, would read on claim 28. In the absence of evidence to the contrary the disclosed prior art streptococcal protein ( $\alpha_2$  M-binding protein) comprise amino acid sequence of SEQ.ID.NO: 6. Since the Office does not have the facilities for examining and comparing applicants' claimed protein with the protein of the prior art, the burden is on applicant to show a novel or unobvious difference between the claimed protein and the protein of the prior art. See *In re Best*, 562 F.2d 1252, 195 USPQ 430 (CCPA 1977).

Applicant's arguments filed on 6/23/03 have been fully considered but they are not deemed to be persuasive.

Applicant states that the cited reference did not disclose the sequence, SEQ.ID.NO: 6 or variant.

The examiner has rejected the claims based on inherency since streptococcal protein ( $\alpha_2$  M-binding protein) bound to the  $\alpha_2$  macroglobulin and therefore, it contains the amino acid sequence SEQ.ID.NO: 6. Applicant did not show any evidence that the disclosed protein and the claimed invention are not the same.

#### ***Status of Claims***

9. No claims are allowed.

#### **Conclusion**

10. **THIS ACTION IS MADE FINAL.** Applicant is reminded of the extension of time policy as set forth in 37 CFR 1.136(a).

A shortened statutory period for response to this final action is set to expire THREE MONTHS from the date of this action. In the event a first response is filed within TWO MONTHS of the mailing date of this final action and the advisory action is not mailed until after the end of the THREE-MONTH shortened statutory period, then the shortened statutory period will expire on the date the advisory action is mailed, and any extension fee pursuant to 37 CFR 1.136(a) will be calculated from the mailing date of the advisory action. In no event will the statutory period for response expire later than SIX MONTHS from the date of this final action.

11. Any inquiry concerning this communication or earlier communications from the examiner should be directed to Padma Baskar whose telephone number is (703) 308-8886. The examiner can normally be reached on Monday through Friday from 6:30 AM to 4 PM EST

If attempts to reach the examiner by telephone are unsuccessful, the examiner's supervisor, Lynette Smith, can be reached on (703) 308-3909. The fax phone number for the organization where this application or proceeding is assigned is (703) 308-4242.

Any inquiry of a general nature or relating to the status of this application or proceeding should be directed to the receptionist whose telephone number is (703) 308-1235.

Padma Baskar Ph.D.

9/16/03

hj  
LYNETTE R. F. SMITH  
SUPERVISORY PATENT EXAMINER  
TECHNOLOGY CENTER 1600